

SEQUENCE LISTING

<110> ASAHIKASEI KOGYO KABUSHIKI KAISHA  
ASAHI MEDICAL CO., LTD.

<120> Separating apparatus of cells and separating method

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85 90 95			
TCC TCC AAC ACA GCC TAC ATG CAG CTC AGC AGC CTG ACA TCT GAG GAC			336
Ser Ser Asn Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu Asp			
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Ser Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Val Leu Thr			
145 150 155 160			
CAA TCT CCA GCT TCT TTG GCT GTG TCT CTA GGG CAG AGG GCC ACC ATC			528
Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly Gln Arg Ala Thr Ile			
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Ser Cys Lys Ala Ser Gln Ser Val Asp Tyr Asp Gly Asp Ser Tyr Met			
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Asn Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr			
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85	90	95	
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195 200 205	
TTA CAT TGG TAC CTG CAG AAG CCA GGC CAG TCT CCA AAT CTC CTG ATC	672
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TAC AAA GTT TCC AAC CGA TTT TCT GGG GTC CCA GAC AGG TTC AGT GGC	720
Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly	
225 230 235 240	
AGT GGA TCA GGG ACA GAA TTC ACA CTC AAG ATC AGC AGA GTG GAG GCT	768
Ser Gly Ser Gly Thr Glu Phe Thr Leu Lys Ile Ser Arg Val Glu Ala	
245 250 255	
GAG GAT CTG GGA GTT TAT TTC TGC TCT CAA AGT ACA CAT GTT CCG CTC	816
Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Leu	
260 265 270	
ACG TTC GGT GCT GGG ACC AAG GTG GAG CTG AAA CGG GCG GCC GCA GGT	864
Thr Phe Gly Ala Gly Thr Lys Val Glu Leu Lys Arg Ala Ala Ala Gly	
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Ala Pro Val Pro Tyr Pro Asp Pro Leu Glu Pro Arg Ala Ala	
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Asn Val Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr	
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GCG GCC CAG CCG GCC ATG GCC CAG GTG AAG CTG CAG CAG TCT GGA CCT	144
Ala Ala Gln Pro Ala Met Ala Gln Val Lys Leu Gln Gln Ser Gly Pro	
35 40 45	
GGC CTA GTG CAG CCC TCA CAG AGC CTG TCC TTC ATC TGC ACA GTC TCT	192
Gly Leu Val Gln Pro Ser Gln Ser Leu Ser Phe Ile Cys Thr Val Ser	
50 55 60	
GGT TTC TCA TTA ACT AGT CAT GGT GTA CAC TGG GTT CGC CAG TCT CCA	240
Gly Phe Ser Leu Thr Ser His Gly Val His Trp Val Arg Gln Ser Pro	
65 70 75 80	
GGA AAG GGT CTG GAG TGG CTG GGA GTG ATA TGG GGT GCT GGA AGG ACA	288
Gly Lys Gly Leu Glu Trp Leu Gly Val Ile Trp Gly Ala Gly Arg Thr	
85 90 95	
GAC TAT AAT GCA GCT TTC ATA TCC AGA CTG AGC ATC AGC AGG GAC ATT	336
Asp Tyr Asn Ala Ala Phe Ile Ser Arg Leu Ser Ile Ser Arg Asp Ile	
100 105 110	
TCC AAG AGC CAA GTT TTC TTT AAG ATG AAC AGT CTG CAA GTT GAT GAC	384
Ser Lys Ser Gln Val Phe Phe Lys Met Asn Ser Leu Gln Val Asp Asp	
115 120 125	
ACA GCC ATA TAT TAC TGT GCC AGA AAT AGG TAC GAG AGC TAC TTT GAC	432
Thr Ala Ile Tyr Tyr Cys Ala Arg Asn Arg Tyr Glu Ser Tyr Phe Asp	
130 135 140	
TAC TGG GGC CAA GGG ACC ACG GTC ACC GTC TCC TCA GGT GGA GGC GGT	480
Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly	
145 150 155 160	
TCA GGC GGA GGT GGC TCT GGC GGT GGC GGA TCG GAC ATC GAG CTC ACT	528
Ser Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Glu Leu Thr	
165 170 175	
CAG TCT CCA CTC TCC CTG CCT GTC AGT CTT GGA GAT CAG GCC TCC ATC	576
Gln Ser Pro Leu Ser Leu Pro Val Ser Leu Gly Asp Gln Ala Ser Ile	
180 185 190	
TCT TGC AGA TCT AGT CAG AAC CTT GTA CAC AGT AAT GGA AAT ACC TAT	624
Ser Cys Arg Ser Ser Gln Asn Leu Val His Ser Asn Gly Asn Thr Tyr	
195 200 205	
TTA CAT TGG TAC CTG CAG AAG CCA GGC CAG TCT CCA AAT CTC CTG ATC	672
Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Asn Leu Leu Ile	
210 215 220	
TAC AAA GTT TCC AAC CGA TTT TCT GGG GTC CCA GAC AGG TTC AGT GGC	720
Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly	
225 230 235 240	
AGT GGA TCA GGG ACA GAA TTC ACA CTC AAG ATC AGC AGA GTG GAG GCT	768
Ser Gly Ser Gly Thr Glu Phe Thr Leu Lys Ile Ser Arg Val Glu Ala	
245 250 255	
GAG GAT CTG GGA GTT TAT TTC TGC TCT CAA AGT ACA CAT GTT CCG CTC	816
Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Leu	
260 265 270	
ACG TTC GGT GCT GGG ACC AAG GTG GAG CTG AAA CGG GCG GCC GCA GGT	864

Thr Phe Gly Ala Gly Thr Lys Val Glu Leu Lys Arg Ala Ala Ala Gly  
275 280 285  
GCG CCG GTG CCG TAT CCG GAT CCG CTG GAA CCG CGT GCC GCA AAG 909  
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